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Search information block:
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Query length: 600
Database: Issued_Patents_AA.*
Database sequences: 212252
Database length: 22503292
Search time (sec): 88.450000

score_list:

seq_documentation block:

Patent No. 6015448

GENERAL INFORMATION:
   APPLICANT: Boyle, William J.
   APPLICANT: Lacey, David L.
   APPLICANT: Calzone, Frank J.
   APPLICANT: Chang, Ming Shi
   TITLE OF INVENTION: OSTEOPROTEGERIN
   NUMBER OF SEQUENCES: 53
   CORRESPONDENCE ADDRESS:
      ADDRESSEE: Amgen Inc.
      STREET: 1840 DeHavilland Drive
      CITY: Thousand Oaks
      STATE: California
      COUNTRY: USA
      ZIP: 91320-1789
   COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: PatentIn Release #1.0, Version #1.30
   CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/704,774,022
      FILING DATE: 12-DEC-1995
      CLASSIFICATION:
   PRIOR APPLICATION DATA:
      APPLICATION NUMBER: US 2,577,788
      FILING DATE:
   ATTORNEY/AGENT INFORMATION:
      NAME: Witex, Robert R.
      REFERENCE/DOCKET NUMBER: A-378
   INFORMATION FOR SEQ ID NO: 46:
   SEQUENCE CHARACTERISTICS:
      LENGTH: 280 amino acids
      TYPE: amino acid
      STRANDEDNESS: single
      TOPOLOGY: linear
   MOLECULE TYPE: protein

alignment_scores:

aligned block:

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;  INFORMATION FOR SEQ ID NO:    46:
;    SEQUENCE CHARACTERISTICS:
;       LENGTH:  280 amino acids
;       TYPE:  amino acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
;    MOLECULE TYPE:  protein
US-08-795-445A-46


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seq_documentation_block:
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; Patent No. 6284485
; GENERAL INFORMATION:
;   APPLICANT: Boyle, Willaim J.
;   APPLICANT: Lacey, David L
;   APPLICANT: Calzone, Frank J.
;   APPLICANT: Chang, Ming-Shi
;   TITLE OF INVENTION: OSTEOPROTEGERIN
;   NUMBER OF SEQUENCES: 53
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Amgen Inc.
;   STREET: 1840 Dehavilland Drive
;   CITY: Thousand Oaks
;   STATE: California
;   COUNTRY: USA
;   ZIP: 91320-1789
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/795,445A
;   FILING DATE:
;   CLASSIFICATION:
;   PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: 08/577,788
;   FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;   NAME: Winter, Robert R.
;   REFERENCE/DOCKET NUMBER: A-378

seq_documentation_block:
; Sequence 46, Application US/08745447A
; Patent No. 6284728
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank T
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: Osteoprotegerin
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: One Amgen Center Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91362-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,447A
; FILING DATE:
; CLASSIFICATION: 514
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A 173C
; INFORMATION FOR SEQ ID NO: 46:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 280 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-795-447A-46

alignment_scores:
Quality: 1117.50   Length: 211
Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

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218 GTGAGAGCGGCTCCTTTCACGCCTTCAGAAAACCACCTCAGACACTGCCTC 267
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; Patent No. 6284740
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David L.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN
; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 DeHavilland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91420-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.40
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/474,186
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A 178
; INFORMATION FOR SEQ ID NO: 46:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 280 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-474-186-46

alignment_scores:
    Quality: 1117.50   Length: 211
    Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

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seq_documentation_block:
; Sequence 46, Application US/08/795446B
; Patent No. 6288032
; GENERAL INFORMATION:
; APPLICANT: Boyle, William J.
; APPLICANT: Lacey, David J.
; APPLICANT: Calzone, Frank J.
; APPLICANT: Chang, Ming-Shi
; TITLE OF INVENTION: OSTEOPROTEGERIN

; NUMBER OF SEQUENCES: 53
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Amgen Inc.
; STREET: 1840 Dehaviland Drive
; CITY: Thousand Oaks
; STATE: California
; COUNTRY: USA
; ZIP: 91320-1789
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,446B
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/577,788
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Winter, Robert B.
; REFERENCE/DOCKET NUMBER: A-378
; INFORMATION FOR SEQ ID NO: 46:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 280 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-795-446B-46

alignment_scores:
    Quality: 1117.50   Length: 211
    Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

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 88 ..........CATAGTGTGTGCCCAAAGGAAAATATATC  117
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seq_documentation_block:
; Sequence 25, Application US/08050319B
; Patent No. 5633145
; GENERAL INFORMATION:
;   APPLICANT: M Feldmann, P W. Gray,
;   APPLICANT: M.J.C. Turner, F.M Brennan
;   TITLE OF INVENTION: Modified human TNFalpha (Tumor
;   TITLE OF INVENTION: Necrosis Factor alpha) Receptor
;   NUMBER OF SEQUENCES: 57
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Reed & Robbins
;   STREET: 635 Bryant Street
;   CITY: Palo Alto
;   STATE: California
;   COUNTRY: USA
;   ZIP: 94301
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1 0, version #1.25
;   CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/050 319B
;   FILING DATE: 10-May-1993
;   CLASSIFICATION: 435
;   ATTORNEY/AGENT INFORMATION:
;   NAME: Robbins, Roberta L
;   REGISTRATION NUMBER: 33,208
;   REFERENCE/DOCKET NUMREP: 5150-0030
;   TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (415) 617-8900
;   TELEFAX: (415) 327-3231
; INFORMATION FOR SEQ ID NO: 25:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 455 amino acids
;   TYPE: amino acid
;   TOPOLOGY: linear
;   MOLECULE TYPE: protein
US-08-050-319B-25

alignment_scores:
  Quality: 1117.50   Length: 211
  Ratio: 5.588   Gaps: 1
  Percent Similarity: 94 787   Percent Identity: 94 787

alignment_block:
US-09-525-998A-11 x US-08-050-319B-25

seq_name: (seq1_f/predata/?/lai/US_COMP rep US-08-050-319B-25

seq_documentation_block:
; Sequence 2, Application US/08021668
; Patent No. 5665859
; GENERAL INFORMATION:
;   APPLICANT: WALLACH, David
;   APPLICANT: BRAKIBUSCH, Cord
;   APPLICANT: VARFLOMEEV, Eugene
;   APPLICANT: BAIKIN, Michael
;   TITLE OF INVENTION: MOLECULES INFLUENCING THE SHEDDING OF
;   TITLE OF INVENTION: THE TNF RECEPTORS, THEIR PREPARATION AND THEIR USE
;   NUMBER OF SEQUENCES: 42
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: BROWDY AND NEIMARK
;   STREET: 419 Seventh Street, N.W., Suite 300
;   CITY: Washington
;   STATE: D.C.
;   COUNTRY: USA
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;     ZIP:  20004
;  COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Floppy disk
;     COMPUTER:  IBM PC compatible
;     OPERATING SYSTEM:  PC-DOS/MS-DOS
;     SOFTWARE:  PatentIn Release #1.0, Version #1.30
;  CURRENT APPLICATION DATA:
;     APPLICATION NUMBER:  US/08/321,668
;     FILING DATE:  12-OCT-1994
;     CLASSIFICATION:  435
;  PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  IL 107268
;     FILING DATE:  12-OCT-1993
;  ATTORNEY/AGENT INFORMATION:
;     NAME:  BROWDY, Roger L.
;     REGISTRATION NUMBER:  25,618
;     REFERENCE/DOCKET NUMBER:  WALLACH-13
;  TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  202-737-3528
;     TELEFAX:  202-737-3528
;     TELEX:  248633
;  INFORMATION FOR SEQ ID NO:  2:
;  SEQUENCE CHARACTERISTICS:
;     LENGTH:  455 amino acids
;     TYPE:  amino acid
;     TOPOLOGY:  linear
;     MOLECULE TYPE:  protein
US-08-321-668-2
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alignment_scores:
              Quality: 1117.50   Length: 211
              Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

alignment_block:
US-09-525-998A-11 x US-08-321-668-2

Align seq 1/1 to: US-08-321-668-2 from: 1 to: 455

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 51 GTGTTGGTGGGAGCAAGCCTCTAGGGGTTATTGGA..........   87
    |||||||||||||||||||||||||||||||||||
 17 uLeuLeuValGlyLeuTyrProSerGlyValIleGlyLeuValProHisL   34

 88 ......    GATAGTGTGTGCCCCAAGGAAAATATATC  117
                 ||||||||||||||||||||||||||||||
 34 euGlyAspArgGluLysCysArgSerValCysProGlnGlyLysTyrIle   50

118 CACTGTAAAATAATTGATTGTGTATGTAAGTGCATAAAGAAGCTA  167
    |||||||||||||||||||||||||||||||||||||||||||||
 51 HisProGlnAsnAsnSerIleCysCysThrLysCysHisLysGlyThrTy   67

168 CTTGTAAATGCATCTGGATGGGGGAGATACGAGATCTAGGGAGT  217
    |||||||||||||||||||||||||||||||||||||||||||
 67 rLeuTyrAsnAspCysProGlyProGlyGlnAspThrAspCysArgGluC   84

216 CTCGCACATGGGTAGGGATGGGGAGAAAGAGGCTCTCTAGGGGC  267
    ||||||||||||||||||||||||||||||||||||||||||||
 84 ysGluSerGlySerPheThrAlaSerGluAsnHisLeuArgHisCysLeu  100

268 AGTCGTGTGCAAATGGGAAAAGGGATGGTGGAGATCTGTCTTGTG  317
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101 SerCysSerLysCysArgLysGlyGluMetGlyValGlnLeuSerSerCy  117

318 CACAGTGACGGGACATGGTGGCCAAGGGGGTGAGGAGAACCACGGC  367
    |||||||||||||||||||||||||||||||||||||||||||||
117 sThrValAspArgAspThrValCysGlyCysArgLysAsnGlnTyrArgH  134

368 ATATTGGAGTGGAAAACCTTTTCCAGTCTTTCCAGTCGGGGTGG  417
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seq_name: /cgn2_6/ptodata/2/i34/5A_COMB.pep US-08-837-941-2

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134 AATGGGAACTTTGGTTGGAGGAAAGAACAGAACACGGCGGAC  467
    ||||||||||||||||||||||||||||||||||||||||||
151 AsnGlyThrValHisLeuSerCysGlnGluLysGliAsnThrValCysH  167

468 CTGCCATGCAGGTTTCTTCTAAGAGAAGAAAACGAGGTGTCTCCTGTAGTA  517
    |||||||||||||||||||||||||||||||||||||||||||||||||||
167 rCysHisAlaGlyPhePheLeuArgGluAsnGluCysValSerCysSerA  184

518 ACTCGAAGAAAAGCCTGGAGTCGCACGAAGTCTGCTGCCTACCCAGATCAG  567
    |||||||||||||||||||||||||||||||||||||||||||||||||||
184 snCysLysCysSerLeuGluCysThrLysLeuCysLeuProGlnIleGlu  200

568 AATGTTAAGGCACTGAGGACTCAGGATCAGGGACTACA  600
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201 AsnValLysGlyThrGluAspGlnThrThr  211
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alignment_scores:
              Quality: 1117.50   Length: 211
              Ratio: 5.588   Gaps: 1

Thu Apr 25 14:25:57 2002

Percent Similarity: 94.787   Percent Identity: 94.787

alignment_block:
US-09-525-998A-11 x US-08-837-941-2 ...

Align seq 1/1 to US 08 837 941 2 from 1 to 455

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   51 GCTGTTGGTGGGAATATACCCTCAGGGGTTATTGGA..............  87
      ||||||||||||||||||||||||||||||||||||
   17 uLeuLeuValGlyIleTyrProGlnGlyLeuValProHisL  34

   88 ..............GATAGTGTGTGTCCCAGGAAAATATATC  117
                  ||||||||||||||||||||||
   34 cuGlyAspArgGlyLysArgAspSerValCysProGlnGlyTyrTyrIle  50

  118 CACCTCAAAATAATTCGATTGCGTGTACCAAGTGCCACAAAGGAAGGACCCA  167
      |||||||||||||||||||||||||||||||||||||||||||||||||||
   51 HisProGlnAsnAsnSerIleCysTrpTyrSerHisTyrSerGlyThrT  67

  168 CTTGTACAATGACTGTCCAGGTGGGCAGGATAGGGACTGCAGGGACT  217
      |||||||||||||||||||||||||||||||||||||||||||||||
   67 rLeuTyrAsnAspCysProGlyProGlnAspArgCysArgGluC  81

  218 GTGAGAGCGGTGTTCTAGAAAACCACCAGACACTGGCTC  267
      |||||||||||||||||||||||||||||||||||||||
   84 ysGluSerGlySerPheThrAlaSerGlnHisLeuArgHisCysLeu  100

  268 AGGCTGCTCCAAATGCCGAAAGGCAAATGGGTCAGGTGGAGATCTCTTG  317
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  101 SerCysSerGlySerArgLysGlyMetGlyGlnValGluIleSerSerCy  117

  318 CACAGTGGCACCGGACACCGGGTGTCCCAGCAAGAACCAGTAGTACCGGC  367
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  368 AATTATTGGAGGAAAACTTTCCAGTGCTTCAATTGGAGACTTGGCTC  417
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  134 IsIleTyrTrpSerGluAsnLeuGlnPheGlnCysPheAsnCysSerLeu  150

  418 AAGGGACCCTCTCTCCCCACCTCCCCCACAAAACACAAAACAGCGTGCCAG  467
      |||||||||||||||||||||||||||||||||||||||||||||||||
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  167 rCysHisAlaGlyPhePheLeuArgGluAsnGluCysAsnGluCysSerA  184

  518 ACTGTAAGAAAAGACTGGAGTGACTACCCTACGCCGGTACCCCATTCAG  567
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  201 AsnValLysGlyThrGluAspSerGlyThrThr  211
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seq_name: /gr2:/gttydata/gexgtech_temp_ff9 US 09 126-016-2

seq_documentation_block:
; Sequence 2, Application US/08126016
; Patent No. 5811261
; GENERAL INFORMATION:
;   APPLICANT: WALLACH, DAVID
;   APPLICANT: NOPHAR, YARON
;   APPLICANT: KIMBER, OLIVER
;   APPLICANT: ENGELMANN, HARTMUT
;   APPLICANT: BRAKEBUSCH, CORD
;   APPLICANT: ADERKA, DAN
;   TITLE OF INVENTION: EXPRESSION OF THE RECOMBINANT TUMOR
;   TITLE OF INVENTION: NECROSIS FACTOR BINDING PROTEIN I (THP-1)

Percent Similarity: 94.787   Percent Identity: 94.787

alignment_scores:
    Quality: 1117.50   Length: 211
    Ratio: 5.588   Gaps: 1
    Percent Similarity: 94.787   Percent Identity: 94.787

alignment_block:
US-09-525-998A-11 x US-08-126-016-2 ...

Align seq 1/1 to US-08-126-016-2 from 1 to 455

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      ||||||||||||||||||||||||||||||||||||
   17 uLeuLeuValGlyIleTyrProGlnGlyLeuValProHisL  34

   88 ..............GATAGTGTGTGTCCCAGGAAAATATATC  117
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   34 euGlyAspArgGlyLysArgAspSerValCysProGlnGlyTyrTyrIle  50

  118 CACCTCAAAATAATTCGATTGCGTGTACCAAGTGCCACAAAGGAAGGACCCA  167
      |||||||||||||||||||||||||||||||||||||||||||||||||||
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  168 CTTGTACAATGACTGTCCAGGTGGGCAGGATAGGGACTGCAGGGACT  217
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  268 AGGCTGCTCCAAATGCCGAAAGGCAAATGGGTCAGGTGGAGATCTCTTG  317
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                                    Quality: 1117.50              Length:  211
                                    Ratio:    5.588               Gaps:      1
Percent Similarity: 94.787          Percent identity: 94.787

alignment_block:
US-09-525-998A-11 x US-08-465-982-25

Align seq 1/1  to: US-08-465-982-25  from: 1  to: 455

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  17 uLeuLeuValGlyIleTyrProSerGlyValIleGlyLeuValProHisL  34

  88 ..................GATAGTGTGTCCCCAAGGAAAATATATC 117
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  34 euGlyAspArgGluLysArgAspSerValCysProGlnGlyLysTyrIle  50

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 218 GTGAAGGGTCTTCTAACTTCAAAAGTCATGATAATGTCTG 267
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 268 AGTGGTGCAAATGCGAAAGGAAATGGGCAGGTGGAGATCTTCTTCTG 317
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 101 SerCysSerLysArgLysGluGlnGlnMetGlyGlnValGlnIleSerSerCy 117

 318 CACAGTGGAGAAACGTGTGGTGTAGAAGAACAGTACAAGGC 367
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 117 sThrValAspArgAspThrValCysGlyCysArgLysAsnGlnTyrArgH 134

 368 ATTATTGGAGTGAAAACCTTTCCAGTGCTTCAATTGCAGCCTCTGCTC 417
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 134 isTyrTrpSerGluAsnLeuPheGlnCysPheAsnCysSerLeuCysLeu 150

 418 AATGGGACCGTGCCACCTCTCCTGCCCAGGAGAAACAGAACACCGTGCAC 467
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 151 AsnGlyThrValHisLeuSerCysGlnGluLysGlnAsnThrValCysTh 167

 468 CTGCCATGCAGTTCTTTCTAAGAGAAAACGATGCTCCTCCTGTAGTA 517
     ||||||||||||||||||||||||||||||||||||||||||||||
 167 rCysHisAlaGlyPheLeuArgGluAsnGluCysValSerCysSerA 184

 518 ACTGTAAGAAACGCTGAGAGTGGCGAAGTTGCCTACCCAGATTGAG 567
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 184 snCysLysLysSerLeuGluCysThrLysSerLeuProGlnIleGlu 200

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 201 AsnValLysGlyThrGluAspSerGlyThrThr 211
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seq_name: /cgmb_2_f/protdata/2/2094/SW_COMB.pep::US-08-465-982-25

seq_documentation_block:
; Sequence 25, Application US/08465982
; Patent No. 5863786
; GENERAL INFORMATION:
;   APPLICANT: M.Feldmann, P.W. Gray,
;   APPLICANT: M.J.C. Turner, F.M Brennan
;   TITLE OF INVENTION: Modified human TNFalpha (Tumor
;   TITLE OF INVENTION: Necrosis Factor alpha) Receptor
;   NUMBER OF SEQUENCES: 57
;   CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Reed & Robbins
;   STREET: 635 Bryant Street
;   CITY: Palo Alto
;   STATE: California
;   COUNTRY: USA
;   ZIP: 94301
;   COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, version #1.25
;   CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08/465,982
;   FILING DATE:
;   CLASSIFICATION:
;   PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US/08/050,319
;   FILING DATE: 10-May-1994
;   ATTORNEY/AGENT INFORMATION:
;   NAME: Robbins, Roberta L.
;   REGISTRATION NUMBER: 33,208
;   REFERENCE/DOCKET NUMBER: 5166-0049
;   TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (415) 617-8999
;   TELEFAX: (415) 327-3231
; INFORMATION FOR SEQ ID NO: 25:
;   SEQUENCE CHARACTERISTICS:
;   LENGTH: 455 amino acids
;   TYPE: amino acid
;   TOPOLOGY: linear
;   MOLECULE TYPE: protein
US-08-465-982-25

alignment_scores.

seq_name: /cgmb_2_f/protdata/2/2094/SW_COMB.pep::US-08-815-469-5

seq_documentation_block:
; Sequence 5, Application US/08815469
; Patent No. 6153402
; GENERAL INFORMATION:
;   APPLICANT: Yu, Guo-Liang
;   APPLICANT: Ni, Jian
;   APPLICANT: Dixit, Vishva
;   APPLICANT: Gentz, Reiner L.
;   APPLICANT: Dillon, Patrick J.
;   TITLE OF INVENTION: Death Domain Containing Receptors

NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
  ADDRESSEE: Sterne, Kessler, Goldstein & Fox, P.L.L.C.
  STREET: 1100 New York Ave., NW, Suite 600
  CITY: Washington
  STATE: DC
  COUNTRY: USA
  ZIP: 20005-3934
COMPUTER READABLE FORM:
  MEDIUM TYPE: Floppy disk
  COMPUTER: IBM PC compatible
  OPERATING SYSTEM: PC-DOS/MS-DOS
  SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
  APPLICATION NUMBER: US/08/815,469
  FILING DATE: HEREWITH
  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: No. 6153402 Yet Assigned
  FILING DATE: 06-FEB-1997
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: US 60/028,711
  FILING DATE: 17-OCT-1996
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: US 60/013,285
  FILING DATE: 12-MAR-1996
ATTORNEY/AGENT INFORMATION:
  NAME: Steffe, Eric K.
  REGISTRATION NUMBER: 36,688
  REFERENCE/DOCKET NUMBER: 1488 0310002/PKS/RPM
TELECOMMUNICATION INFORMATION:
  TELEPHONE: 202-371-2600
  TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 5:
  SEQUENCE CHARACTERISTICS:
  LENGTH: 455 amino acids
  TYPE: amino acid
  STRANDEDNESS: not relevant
  TOPOLOGY: not relevant
  MOLECULE TYPE: protein
US-08-815-469-5

alignment_scores:
  Quality: 1117.50   Length: 211
  Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

alignment_block:
US-09-525-998A-11 x US-08-815-469-5

Align seq 1/1 to: US-08-815-469-5 from: 1 to: 455

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    ||||||||||||||||||||||||||||||||||||||||
 17 uLeuLeuValGlyIleTyrProSerGlyValIleGlyValLeuProHisL 34

 88 .........GATA...GTGTGTGCCCCAAGGAAAATATATC 117
    ||||||||||||||||||||||||||||||||||
 34 euGlyAspArgGlyLysArgAspSerValCysProValCysProGlySerPhe 50

118 CACCTCAAAATAATTCGATTTGCTGTACCAAGTGCA...AAAGGAACCTA 167
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 51 HisProGlnAsnAsnSerIleCysThrLysLys...GlyThrTy 67

168 CTTGTACAATGACTGTCCAGTCCCCGGGCCAGGATACGGACTGCAGGGAGT 217
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seq_documentation_block:
  Sequence 3, Application US/09006353A
  Patent No. 6261801
GENERAL INFORMATION:
  APPLICANT: WEI, YING-FEI
  APPLICANT: YU, GUO-LIANG
  APPLICANT: GENTZ, REINER
  APPLICANT: RUBEN, STEVEN
  TITLE OF INVENTION: TUMOR NECROSIS FACTOR RECEPTOR 5
  NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
  ADDRESSEE: HUMAN GENOME SCIENCES, INC.
  STREET: 9410 KEY WEST AVENUE
  CITY: ROCKVILLE
  STATE: MD
  COUNTRY: US
  ZIP: 20850
COMPUTER READABLE FORM:
  MEDIUM TYPE: Floppy disk
  COMPUTER: IBM PC compatible
  OPERATING SYSTEM: PC-DOS/MS-DOS
  SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
  APPLICATION NUMBER: 09/006,353A
  FILING DATE:
  CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
  NAME: BROOKES, ANDERS A.
  REGISTRATION NUMBER: 36,373
  REFERENCE/DOCKET NUMBER: PF145
TELECOMMUNICATION INFORMATION:
  TELEPHONE: (301) 309-8504
  TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 3:
  SEQUENCE CHARACTERISTICS:
  LENGTH: 455 amino acids
  TYPE: amino acid
  STRANDEDNESS: single
  TOPOLOGY: linear

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; MOLECULE TYPE: protein
US-09-006-353A-4

alignment_scores:
        Quality: 1117.50   Length: 211
          Ratio: 5.588   Gaps: 1
Percent Similarity: 94.787   Percent Identity: 94.787

alignment_block:
US-09-525-998A-11 x US-09-006-353A-4 ..

Align seq 1/1 to: US-09-006-353A-4   from: 1 to: 455

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   51 GCTGTTGGGGGAATATACCCTCAGGGGTTATTGGA.............. 87
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   17 uLeuLeuValGlyIleTyrProSerGlyValIleGlyValLeuValProHisL 34

   88 .........GATAGTGTGTGCCCCAAGAAAAATATATC 117
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   34 euGlyAspArgGluLysArgSerValCysProGlyGlyCysTyrIle 50

  118 CACCCTCAAAATAATTGGATTTGCTGTACAAGTGCCAAAGGAACCTA 167
      ||||||||||||||||||||||||||||||||||||||||||||||
   51 HisProGlnAsnAsnSerIleCysCysThrLysCysGlnThrTyr 67

  168 CTTGTACAAATGACTGTCCAGGCCCGGGCAGGATACCGACTGCAGGGAGT 217
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   67 rLeuTyrAsnAspCysProGlyProGlyGlnAspThrAspCysArgGluC 84

  218 GTGAGAGCGGTCTTCAGGTTCAGAAAAAAGCTCAGAACTGCCTC 267
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  268 AGTTGTTCCAAATGCCAAAGGGAAAGTGGTCAGGTGGAGATCTTCTTG 317
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  101 SerCysSerLysCysGlnArgLysGluLeuGlyGlnValGluIleSerSerCy 117

  318 CACAGTGACCGGGACACCGCGTGTCGCTGCGAGGAGAACCAGTACCGGC 367
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  117 sThrValAspArgAspThrValCysGlyCysArgLysAsnGlnTyrArgH 134

  368 ATTATTGGAGTGAAAACCTTTCAGTAGTTCAATTGAGAGCCTCGACTC 417
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  418 AATGGGAACGTCAACTCTCCTGGCAGGAGAAACAGAACACATGTGCAC 467
      ||||||||||||||||||||||||||||||||||||||||||||||
  151 AsnGlyThrValAsnSerLeuSerGlyArgGluLysGlnAsnThrValCysTh 167

  468 CTGCCATGCAGGTTTCTTCTCAAGAAGAAAACGAGTGTGTCCCTGTAGTA 517
      ||||||||||||||||||||||||||||||||||||||||||||||||||
  167 rCysHisAlaGlyPhePheLeuArgLysAsnGluCysValSerCysSerA 184

  518 ACTGTAAGAAAAGCTGAGTGCAGAAGTGTGCCTACCCCAGATTGAG 567
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  184 SerCysLysLysSerGluCysThrTyrLysLeuProToLeuProGlu 200

  568 AATGGTTAAGGCACTGAGGACTCAGCACCACTA 600
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  201 AsnValLeuGlyThrGluAspSerGlnThrThr 211
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seq_document.block:
seq_name: /vgn2_4/predata/2/baa/5B_COMB_pep:US-09-086-483A-5
; Sequence 5, Application US/09086483A
; Patent No. 6214580
; GENERAL INFORMATION:
; APPLICANT: NI, et al.

alignment_scores:
          Quality: 1049.50   Length: 199
          Ratio: 5.612   Gaps: 1
     Percent Similarity: 93.970   Percent Identity: 93.970

alignment_block:
     us-09-525-998A-11 x us-08-050-419R-48

Align seq 1/1 to 35 08 050 419R-48 from 1 to 199

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 51 GCTGTTCGTGGAAGTACCCGCCACGGCTATTGCA........  87
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 17 AlaLeuValPhePheProGlyValIleGlnLeuValProHisL  34

 88 ........GATAGTGTGTGCAAGGAAAAATATC  117
            |||||||||||||||||||||||
 34 CysGlyAspArgCysLysSerLysLysLysTyrTyrPhe  50

118 CAGGTCAAAAATGTGTCTGTAGAAGTGCACCTA  167
    ||||||||||||||||||||||||||||||||
 51 GlnValLysAsnValSerValGluValHisLeu  67

168 CTTGTACAAGAGTGCAGGGTCAAGTCCAGCC  217
    |||||||||||||||||||||||||||||||
 67 LeuValGlnGluCysArgValLysSerSerA  84

    AAAGTCAAAAACAGAACCA  267
    ||||||||||||||||||
 84 LysGlnHisCysTyrCysLeu  100

268 AGTGCTCCAAATGCGAAGGAAATGGGTCAGTGGATCTTCTTG  317
    ||||||||||||||||||||||||||||||||||||||||
101 SerCysSerLysTyrSerArgCysSerMetGlyGlnValGlu  117

318 CACAGTGGACGGGACACCGTGTGGCTGCAGCAAGAACCAGTACCGGC  367
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117 ThrValAspArgAspThrValCysGlyCysLysLysAsnGlnTyrArgH  134

368 ATTATTGGACGGAAACCTTCCAGTGGTTCAATGCAGCCGTGCCTC  417
    |||||||||||||||||||||||||||||||||||||||||
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418 AATGGGACGGTGACCTCTTGCAGAAGAAACAGAACAGGTGTGGAC  467
    |||||||||||||||||||||||||||||||||||||||||
151 AsnGlyThrValThrSerCysArgGluAsnHisValCysSth  167

468 CTGCATGCAGGTTCTTTTAAGAGAAAACGAGTGTGTGCGCAGTA  517
    |||||||||||||||||||||||||||||||||||||||||
167 CysHisAlaGlyPheLeuLysArgGluAsnGluCysValSerCysSerA  184

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seq_documentation_block:
; Sequence 48, Application US/08050319B
; Patent No. 5633145
; GENERAL INFORMATION:
; APPLICANT: M.Feldmann, P.W. Gray,
; APPLICANT: M.J.C. Turner, F.M Brennan
; TITLE OF INVENTION: Modified human TNFalpha (Tumor
; TITLE OF INVENTION: Necrosis Factor alpha) Receptor
; NUMBER OF SEQUENCES: 57
; CORRESPONDENCE ADDRESS:
;   ADDRESSEE: Reed & Robbins
;   STREET: 635 Bryant Street
;   CITY: Palo Alto
;   STATE: California
;   COUNTRY: USA
;   ZIP: 94301
; COMPUTER READABLE FORM:
;   MEDIUM TYPE: Floppy disk
;   COMPUTER: IBM PC compatible
;   OPERATING SYSTEM: PC-DOS/MS-DOS
;   SOFTWARE: PatentIn Release #1.0, version #1.25
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/08050319R
;   FILING DATE: 10-May-1993
;   CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
;   NAME: Robbins, Robert J
;   REGISTRATION NUMBER: 33,208
;   REFERENCE/DOCKET NUMBER: 5150-0030
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (415) 617-8499
;   TELEFAX: (415) 327-3231
; INFORMATION FOR SEQ ID NO: 48:
; SEQUENCE CHARACTERISTICS:
;   LENGTH: 199 amino acids
;   TYPE: amino acid
;   TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-050-419R-48

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:    STREET:  635 Bryant Street
:    CITY:  Palo Alto
:    STATE:  California
:    COUNTRY:  USA
:    ZIP:  94301
:  COMPUTER READABLE FORM:
:    MEDIUM TYPE:  Floppy disk
:    COMPUTER:  IBM PC compatible
:    OPERATING SYSTEM:  PC-DOS/MS-DOS
:    SOFTWARE:  Patent In Release #1.0, version #1.25
:  CURRENT APPLICATION DATA:
:    APPLICATION NUMBER:  US/08/465,982
:    FILING DATE:
:    CLASSIFICATION:
:  PRIOR APPLICATION DATA:
:    APPLICATION NUMBER:  US/08/464,519
:    FILING DATE:  10-May-1993
:  ATTORNEY/AGENT INFORMATION:
:    NAME:  Robbins, Roberta L.
:    REGISTRATION NUMBER:  33,208
:    REFERENCE/DOCKET NUMBER:  5150-0030
:  TELECOMMUNICATION INFORMATION:
:    TELEPHONE:  (415) 617-8999
:    TELEFAX:  (415) 327-3231
:  INFORMATION FOR SEQ ID NO:  48:
:    SEQUENCE CHARACTERISTICS:
:      LENGTH:  199 amino acids
:      TYPE:  amino acid
:      TOPOLOGY:  linear
:    MOLECULE TYPE:  protein
US-08-465-982-48
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alignment_scores:
            Quality:  1049.50            Length:       199
              Ratio:  5.612                Gaps:         1
Percent Similarity:  93.970    Percent Identity:  93.970

alignment_block:
US-09-525-998A-11  to: US-08-465-982-48  ..

Align seq 1/1  to: US-08-465-982-48  from: 1  to: 199

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  51 GCTGTTGGTGGGGAATATACCCCTCCAGGGGTTATGGGA.........  87
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  17 uLeuLeuValGlyIleTyrProGlyValIleGlyLeuValProHisL  34

  88                GATAGTTGTGTGTCCAAGGAAAATATATC  117
                    |||||||||||||||||||||||||||||
  34 euGlyAspArgCysValArgSerValCysProThrCysGlySerPhe  50

 118 CACCCTCAAAATAATTCGATTGCCATTGCTCGACCAGCCCAAAGGAACCTA  167
     ||||||||||||||||||||||||||||||||||||||||||||||||||
  51 HisProGlnAsnAsnSerIleCysCysThrLysCysHisLysGlyThrTy  67

 168 CTTGTACAATGACTGTCCAGAGTCCCAGGATAACCGGACTGCAGGGAGT  217
     ||||||||||||||||||||||||||||||||||||||||||||||||
  67 rLeuTyrAsnAspCysProGlyProGlnAspThrAspCysArgGluC  84

 218 GTGAGAGCGGCTGCTTCACCGGTTCAGAAAATCACCTCAGACACTGCCTC  267
     |||||||||||||||||||||||||||||||||||||||||||||||||
  84 ysGluSerGlySerPheThrAlaSerGluAsnHisLeuArgHisCysLeu  100

 268 AGCTGCTCCAAAGTGCCCAAAGAAATGGGTCAGTCGAGATCTCTTCTTG  317
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 318 CACAGTGGCACCGGACACCGTGTGTGGCTGCAGAAGAACCAGTACCGGC  367
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us-09-525-998a-11.rai

Thu Apr 25 14:25:57 2002